All samples combined
Methylation ratios sample−mean[controls]
CG.ratio_DE
CHG.ratio_DE
CHH.ratio_DE
CG.DMR
CHG.DMR
CHH.DMR
DMR context
0.20
0.15
0.10
0.05
0.00
−100
−50
0
50
100
Methylation ratios context
Pete's agressive filtering